

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/083,815A

DATE: 08/12/2002 TIME: 14:07:23

Input Set : A:\435c2.app

Output Set: N:\CRF3\08122002\J083815A.raw

```
5 <110> APPLICANT: Anderson, Christen M.
              Clevenger, William
     9 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR REGULATING
              ENDOGENOUS INHIBITOR OF ATP SYNTHASE, INCLUDING
     10
              TREATMENT FOR DIABETES
     11
     14 <130> FILE REFERENCE: 660088.435C2
C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/083,815A
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18 <141> CURRENT FILING DATE: 2002-02-27

20 <160> NUMBER OF SEQ ID NOS: 73

22 <170> SOFTWARE: FastSEQ for Windows Version 4.0

24 <210> SEQ ID NO: 1

25 <211> LENGTH: 6

26 <212> TYPE: PRT

27 <213> ORGANISM: Artificial Sequence

29 <220> FEATURE:

30 <223> OTHER INFORMATION: Epitope tag

32 <400> SEQUENCE: 1

33 His His His His His

36 <210> SEQ ID NO: 2 37 <211> LENGTH: 7

38 <212> TYPE: PRT

39 <213> ORGANISM: Artificial Sequence

41 <220> FEATURE:

42 <223> OTHER IN. ORMATION: Epitope tag

44 <400> SEQUENCE: 2

45 Asp Tyr Asp Asp Asp Lys

46 1

48 <210> SEQ ID NO: 3

49 <211> LENGTH: 6

50 <212> TYPE: PRT

51 <213> ORGANISM: Artificial Sequence

53 <220> FEATURE:

54 <223> OTHER INFORMATION: Epitope tag

56 <400> SEQUENCE: 3

57 Asp Thr Tyr Arg Tyr Ile

58 1

60 <210> SEQ ID NO: 4

61 <211> LENGTH: 6

62 <212> TYPE: PRT

63 <213> ORGANISM: Artificial Sequence

65 <220> FEATURE:

66 <223> OTHER INFORMATION: Epitope tag

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68 <400> SEQUENCE: 4 69 Thr Asp Phe Tyr Leu Lys 72 <210> SEQ ID NO: 5 73 <211> LENGTH: 10 74 <212> TYPE: PRT 75 <213> ORGANISM: Artificial Sequence 77 <220> FEATURE: 78 <223> OTHER INFORMATION: Epitope tag 80 <400> SEQUENCE: 5 81 Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu 82 1 84 <210> SEQ ID NO: 6 85 <211> LENGTH: 9 86 <212> TYPE: PRT 87 <213> ORGANISM: Artificial Sequence 89 <220> FEATURE: 90 <223> OTHER INFORMATION: Epitope tag 92 <400> SEQUENCE: 6 93 Glu Glu Glu Glu Tyr Met Pro Met Glu 94 1 96 <210> SEQ ID NO: 7 97 <211> LENGTH: 9 98 <212> TYPE: PRT 99 <213> ORGANISM: Artificial Sequence 101 <220> FEATURE: 102 <223> OTHER INFORMATION: Epitope tag 104 <400> SEQUENCE: 7 105 Tyr Pro Tyr Asp Val Pro Asp Tyr Ala 106 1 108 <210> SEQ ID NO: 8 109 <211> LENGTH: 5 110 <212> TYPE: PRT 111 <213> ORGANISM: Artificial Sequence 113 <220> FEATURE: 114 <223> OTHER INFORMATION: Epitope tag 116 <400> SEQUENCE: 8 117 Arg Tyr Ile Arg Ser 118 1 120 <210> SEQ ID NO: 9 121 <211> LENGTH: 6 122 <212> TYPE: PRT 123 <213> ORGANISM: Artificial Sequence 125 <220> FEATURE: 126 <223> OTHER INFORMATION: Epitope tag 128 <400> SEQUENCE: 9 129 Pro Pro Glu Pro Glu Thr 130 1 132 <210> SEQ ID NO: 10

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133 <211> LENGTH: 8
134 <212> TYPE: PRT
135 <213> ORGANISM: Artificial Sequence
137 <220> FEATURE:
138 <223> OTHER INFORMATION: cellular transport sequence
140 <400> SEQUENCE: 10
141 Arg Lys Lys Arg Arg Gln Arg Arg
142 1
144 <210> SEQ ID NO: 11
145 <211> LENGTH: 21
146 <212> TYPE: DNA
147 <213> ORGANISM: Artificial Sequence
149 <220> FEATURE:
150 <223> OTHER INFORMATION: cellular transport sequence
152 <400> SEQUENCE: 11
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153 aggaagaagc ggagacagag a
155 <210> SEQ ID NO: 12
156 <211> LENGTH: 324
157 <212> TYPE: DNA
158 <213> ORGANISM: Rattus norvegicus
160 <400> SEQUENCE: 12
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161 atggcaggct cggcgttggc ggttcgggct cggctcggtg tctgggggtat gagggtcctg
162 caaaccegag getteggete ggaetegteg gagageatgg attegggege tggeteeate
                                                                            120
163 cgagaagctg gtggggcctt cgggaaacga gagaaggctg aagaggatcg gtacttccga
                                                                            180
164 gagaagacta gagagcagct ggctgccttg aagaagcacc atgaagatga gattgaccac
                                                                            240
165 cattcgaagg agatagagcg tctgcaaaaa cagatcgaac ggcataagaa gaagattaaa
                                                                            300
                                                                            324
166 tacctaaaga atagtgagca ttga
168 <210> SEQ ID NO: 13
169 <211> LENGTH: 107
170 <212> TYPE: PRT
171 <213> ORGANISM: Rattus norvegicus
173 <400> SEQUENCE: 13
174 Met Ala Gly Ser Ala Leu Ala Val Arg Ala Arg Leu Gly Val Trp Gly
                                         10
175 1
                      5
176 Met Arg Val Leu Gln Thr Arg Gly Phe Gly Ser Asp Ser Ser Glu Ser
177
178 Met Asp Ser Gly Ala Gly Ser Ile Arg Glu Ala Gly Gly Ala Phe Gly
            35
179
180 Lys Arg Glu Lys Ala Glu Glu Asp Arg Tyr Phe Arg Glu Lys Thr Arg
                             55
182 Glu Gln Leu Ala Ala Leu Lys Lys His His Glu Asp Glu Ile Asp His
                                             75
                         70
184 His Ser Lys Glu Ile Glu Arg Leu Gln Lys Gln Ile Glu Arg His Lys
                                         90
185
                     85
186 Lys Lys Ile Lys Tyr Leu Lys Asn Ser Glu His
                100
189 <210> SEQ ID NO: 14
190 <211> LENGTH: 75
191 <212> TYPE: DNA
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192 <213> ORGANISM: Rattus norvegicus
194 <400> SEQUENCE: 14
195 atggcaggct cggcgttggc ggttcgggct cggctcggtg tctggggtat gagggtcctg
                                                                             60
                                                                             75
196 caaacccgag gcttc
198 <210> SEQ ID NO: 15
199 <211> LENGTH: 509
200 <212> TYPE: DNA
201 <213> ORGANISM: Mus musculus
203 <400> SEQUENCE: 15
204 cgcaacgcga gctgagcaac gccgaagaca atggcaggct cggcgttggc agttcgggct
                                                                             60
                                                                            120
205 cqqttcqqtq tctqqqqtat gaaggtcctg caaacccgag gcttcgtctc ggactcgtcg
206 gatagcatgg atacgggcgc tggctccatc cgagaagctg gtggagcctt cggaaaacga
                                                                            180
207 gaaaaggctg aagaggatcg gtacttccga gagaagacta aagaacagct ggctgccctg
                                                                            240
                                                                            300
208 aggaaacacc atgaagatga gattgaccac cattcgaagg agatagagcg tctgcagaag
209 caaattgatc gccataagaa gaagatccaa caactaaaga ataatcattg aatgcgcgca
                                                                            360
                                                                            420
210 gtcqqtccct cacagagtgg cccqtatcac tccccacqtc tgtagacaca tggctttgaa
211 tgattactat ttggtctgtg tgctactaac agataataaa cgatcaccag gaaactttta
                                                                            480
                                                                            509
212 aaaaaaaaa aaaaaaaaa aaaaaaaaa
214 <210> SEQ ID NO: 16
215 <211> LENGTH: 106
216 <212> TYPE: PRT
217 <213> ORGANISM: Mus musculus
219 <400> SEQUENCE: 16
220 Met Ala Gly Ser Ala Leu Ala Val Arg Ala Arg Phe Gly Val Trp Gly
221
                                        10
222 Met Lys Val Leu Gln Thr Arg Gly Phe Val Ser Asp Ser Ser Asp Ser
                20
                                    25
224 Met Asp Thr Gly Ala Gly Ser Ile Arg Glu Ala Gly Gly Ala Phe Gly
                                40
            35
226 Lys Arg Glu Lys Ala Glu Glu Asp Arg Tyr Phe Arg Glu Lys Thr Lys
                            55
                                                 60
228 Glu Gln Leu Ala Ala Leu Arg Lys His His Glu Asp Glu Ile Asp His
                        70
                                             75
229 65
230 His Ser Lys Glu Ile Glu Arg Leu Gln Lys Gln Ile Asp Arg His Lys
                                        90
232 Lys Lys Ile Gln Gln Leu Lys Asn Asn His
                                    105
233
                100
235 <210> SEQ ID NO: 17
236 <211> LENGTH: 23
237 <212> TYPE: DNA
238 <213> ORGANISM: Artificial Sequence
240 <220> FEATURE:
241 <223> OTHER INFORMATION: PCR primer
243 <400> SEQUENCE: 17
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244 cacaaagata tcggaaccct cta
246 <210> SEO ID NO: 18
247 <211> LENGTH: 25
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
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	<220> FEATURE:	
	<223> OTHER INFORMATION: PCR primer <400> SEQUENCE: 18	
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	<211> LENGTH: 47	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: PCR primer	
	<400> SEQUENCE: 19	
	tqaqctcaqa tatqqcagga agaagcggag acagagagga atggcag	47
	<210> SEQ ID NO: 20	
	<211> LENGTH: 34	
270	<212> TYPE: DNA	
271	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
274	<223> OTHER INFORMATION: PCR primer	
276	<400> SEQUENCE: 20	
277	atataagctt tcaatgctca ctattcttta ggta	34
279	<210> SEQ ID NO: 21	
280	<211> LENGTH: 33	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Tat-derived cellular targeting sequence	
	<400> SEQUENCE: 21	•
	agatatggca ggaagaagcg gagacagaga gga	33
	<210> SEQ ID NO: 22	
	<211> LENGTH: 11	
	<212> TYPE: PRT	
	<213> ORGANISM: Artificial Sequence <220> FEATURE:	
	<pre><220> FEATURE: <223> OTHER INFORMATION: Tat-derived cellular targeting sequence</pre>	
	<223> OTHER INFORMATION. Tat delived certain targeting sequence <400> SEQUENCE: 22	
	Arg Tyr Gly Arg Lys Lys Arg Arg Gln Arg Gly	
300	1 5 10	
	<210> SEQ ID NO: 23	
	<211> LENGTH: 48	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
308	<223> OTHER INFORMATION: PCR primer	
	<400> SEQUENCE: 23	
	tgagctcagg atatggcagg aagaagcgga gacagagagg aggctcgg	48
	<210> SEQ ID NO: 24	
314	<211> LENGTH: 34	
	<212> TYPE: DNA	
316	<213> ORGANISM: Artificial Sequence	
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VERIFICATION SUMMARY

DATE: 08/12/2002

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Input Set : A:\435c2.app

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 $L:17\ M:270\ C:$ Current Application Number differs, Wrong Format